# Metabolic network analysis and regulation

Lucas Marmiesse

INRA CNRS, Laboratoire des Interactions Plantes-Microorganismes (LIPM), UMR441, F-31326 Castanet-Tolosan, France

21 novembre 2014





## Metabolism



#### A biochemical reaction

# Metabolism



Example of a metabolic pathway : The Krebs cycle

# Metabolism



## Metabolism regulation



# Metabolism regulation





# **Biological networks**



#### *Ralstonia solanacearum* With Ludovic Cottret and Remi Peyraud



Reconstructed metabolic network.

Reconstructed virulence regulatory network.

Arabidopsis Thaliana defence response Supervisors : Susana Rivas and Philippe Besse

















Constraint based method : Flux Balance Analysis (FBA)



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 Allows to calculate the fluxes (rates) of the reactions of the metabolic network at steady state.



 $\begin{array}{l} \mbox{Production} \\ \mbox{V1+V2} - \mbox{V3} = 0 \ \mbox{mmol/g/h} \\ \mbox{Consumption} \end{array}$ 

Constraint based method : Flux Balance Analysis (FBA)

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R1:	A_ext → A		R1	R2	R3	R4	R5	R6	R7	R8	R9
R2:	$A \rightarrow B$	А	1	-1	-1						
R4 :	$B + E \rightarrow 2D$	В		1		-1		-2			
R5:	E_ext → E	С			1			1	-1		
R6: 87·	2B→C+F C→D	D				2			1	-1	
R8:	$D \rightarrow D_{ext}$	Е				-1	1				
R9:	$F \rightarrow F_{ext}$	F						1			-1

S

http://bio.freelogy.org/wiki/User:JeremyZucker

Constraint based method : Flux Balance Analysis (FBA)

						S							V v1	1
R1:	A_ext → A		R1	R2	R3	R4	R5	R6	R7	R8	R9		v1 v2	
R2:	A → B A → C	Α	1	-1	-1							]	v3	1
R4:	B + E → 2D	В		1		-1		-2					v4	1
R5 :	$E_{ext} \rightarrow E$	с			1			1	-1				v5	1
R6: R7·	2B→C+F C→D	D				2			1	-1		X	v6	1
R8:	D → D_ext	E				-1	1						v7	1
R9:	F → F_ext	F						1			-1		v8	1
													v9	1

Constraint based method : Flux Balance Analysis (FBA)

 Allows to calculate the fluxes (rates) of the reactions of the metabolic network at steady state.



S.v = 0

 $\begin{array}{l} dA/dt = v1 - v2 - v3 = 0 \\ dB/dt = v2 - v4 - 2*v6 = 0 \\ dC/dt = v3 + v6 - v7 = 0 \\ \hline dD/dt = 2*v4 + v7 - v8 = 0 \\ dE/dt = -v4 + v5 = 0 \\ dF/dt = -v4 - v9 = 0 \end{array}$ 

Méthode basée sur des contraintes : Analyse d'équilibre des flux (FBA)

- ► Allows to calculate the fluxes (rates) of the reactions of the metabolic network at steady state.
- Based on the stoichiometry of the network and constraints on the input fluxes. linéaires).
- Resolves a system of linear equations with an objective function.



Orth et al. 2011



### Constraint based methods phylogeny



Lewis et al. 2013

### Metabolic network regulation



## Metabolic network regulation



de Oliveira Dal'Molin et al. 2010

## Metabolic network regulation



de Oliveira Dal'Molin et al. 2010

# Regulatory network

#### Nature

- Composed of biological entities of different types that interact in different ways.
- ► Cause a specialisation (reversible or not) of the metabolism.

#### Analyses and simulations

▶ Many methods (de Jong *et al.* 2002, Karlebach *et al.* 2008).

#### Boolean model

- Each component has only two possible states.
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Multi-state model

- Each component has a finite number of possible states.
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#### Simulations

	Α			В			C	
Condition	on Value		Condition	Value	alue		ndition	Value
B=1	1		A=1	0		A=1	XOR B=1	1
Default	C		Default	1		A=1	AND B=1	2
						D	efault	0
		T0	T1	T2	Т	3	T4	
	А	0						
	В	1						
	С	0						

#### Simulations

4	A			В			C	
Condition	ı Value		Condition	Value	Value		ndition	Value
B=1	1		A=1	0		A=1	XOR B=1	1
Default	C	)	Default	1		A=1	AND B=1	2
						D	efault	0
[		T0	T1	T2	Т	3	T4	
(	А	0	1					
	В	1	1					
	С	0	1					

#### Simulations

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[		T0	T1	T2	Т	3	T4	]
(	А	0	1	1	(	)		
	В	1	1	0	(	)		
	С	0	1	2		1		]

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	I					D	efault	0
[		Т0	T1	T2	Т	3	T4	]
(	А	0	1	1	(	)	0	
	В	1	1	0	(	)	1	
	С	0	1	2		1	0	]



#### Qualitative values

Quantitative values



Constrain the FBA with regulatory network states

► Find a steady state of the regulatory network, constrain the reactions and perform a FBA analysis.

Regulatory network steady state R1 = 0 R2 = 1 R1 = 0C < R2 < C

Constrain the FBA with regulatory network states

- Find a steady state of the regulatory network, constrain the reactions and perform a FBA analysis.
- What network state to use ?

Regulatory network steady state R1 = 0 R2 = 1 R1 = 0C < R2 < C

Constrain the FBA with regulatory network states

- ► Find a steady state of the regulatory network, constrain the reactions and perform a FBA analysis.
- What network state to use ?

	Т0	T1	T2	T3	Τ4
А	0	1	1	0	0
В	1	1	0	0	1
С	0	1	2	1	0

► How to transform a qualitative state into a constraint ?

# Qualitative state to constraint translation



### Qualitative state to constraint translation



### Qualitative state to constraint translation



# What network state to use ?

	Т0	T1	T2	T3	T4
A	0	1	1	0	0
В	1	1	0	0	1
С	0	1	2	1	0

### What network state to use ?

	T0	T1	T2	T3	T4
Α	0	1	1	0	0
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#### What network state to use ?

	Т0	T1	T2	T3	T4
Α	0	1	1	0	0
В	1	1	0	0	1
С	0	1	2	1	0





 $\begin{array}{ll} 0 < C < 0 \\ 0 < C < 5 \\ 5 < C < + inf \\ 0 < C < 5 \end{array} < => 5/4 < C < + inf \\ \end{array}$ 

Dynamic analysis

 Metabolite concentrations allow to go back from FBA results to a regulatory network state.

Logical regulatory network





Dynamic analysis

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Dynamic analysis

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#### Dynamic analysis

 Metabolite concentrations allow to go back from FBA results to a regulatory network state.



# What tools are available for these analyses ?

### File formats

- Metabolic network : Systems Biology Markup Language (SBML). (Hucka *et al.* 2003)
- Regulatory network : SBML qualitative extension. (Chaouiya et al. 2013)

### Software

- ► FBA : Cobra toolbox, surrey FBA, optFlux ...
- ► Boolean regulatory network : Boolnet.
- ► Multi-state regulatory network : GINsim, The Cell Collective.

# FlexFlux

Java software

- Allows analysis of both metabolic and regulatory networks.
- ► Supports SBML and SBML-qual file formats.
- Features steady state and dynamic analysis of the regulatory network.
- ► Features multiple FBA based functions.
- Documentation and examples available at http://lipm-bioinfo.toulouse.inra.fr/flexflux/

800	FlexFlux		
	Choose your analysis :	FBA	Run
Descrip Flexflux Compu	otion : «FBA [options] tes an FBA given a metabolic Required arguments -cond -s	FBA Test DR Pareto CompFVA ERA Reac TwoReacs	<ul> <li>ttion and constraints.</li> <li>Choose file</li> <li>Choose file</li> </ul>
	Optional arguments		
	-senFile		Choose file
	-pre	6	
	-lib	0.0	
	-states		Choose file
	-out		Choose file
	-sol	CPL	EX 💌
	-int		Choose file
	-plot		
	-ext		
	-h		

800	FlexFlux							
	Choose your analysis : FBA	Run						
Description : FlexfluxFBA [options] Computes an FBA given a metabolic network, an objective function and constraints.								
	Required arguments							
	-cond	Choose file						
	-s	Choose file						
	Optional arguments							
	-senFile	Choose file						
	-рге	6						
	-lib	0.0						
	-states	Choose file						
	-out	Choose file						
	-sol	CPLEX 🔻						
	-int	Choose file						
	-plot							
	-ext							
	-h							

😕 🗐 🔲 FBA results								
obj : 0.873922								
Search for an entity :								
Entity name	Value 🔻	٦						
R_ATPS4r	45.51401	•						
R_CYTBD	43.598985							
R_NADH16	38.53461							
R_EX_h2o_e	29.175827							
R_EX_co2_e	22.809833							
R_02t	21.799493							
R_EX_h_e	17.530865							
R_GAPD	16.023526							
R_ENO	14.71614							
R_GLCpts	10.0							
R_PDH	9.282533							
R_ATPM	8.39							
R_PFK	7.477382							
R_FBA	7.477382							
R_TPI	7.477382							
R_ICDHyr	6.00725							
R_ACONTa	6.00725							
R_ACONTb	6.00725							
R_CS	6.00725							
R_FUM	5.064376							
R_SUCDi	5.064376							
R_AKGDH	5.064376							
R_MDH	5.064376							
R_PGL	4.959985							
R_G6PDH2r	4.959985	•						

	Stead	y state	analys	is resul	ts						
Search for an entity :											
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Entity n	1 🔺	2	3	4	5	6	7	8			
KCS16_p	0	0	0	0	0	0	0	0	•		
HCD1	0	0	0	0	1	1	1	1			
KCS3_m	0	0	0	0	0	0	0	0			
KCS5_m	0	0	0	0	0	0	0	0	=		
CER4	0	0	0	0	1	1	1	1			
KCS1_p	0	0	0	0	0	0	1	1			
MIEL4_p3	0	0	0	0	0	0	0	0			
sPLA2_a	0	0	0	0	0	0	0	0			
MIEL4_m	0	0	0	0	0	0	0	0			
MIEL3_m	0	0	0	0	0	0	0	0			
KCS4_p	0	0	0	0	0	0	0	0			
МҮВЗ1_р	0	0	0	0	0	0	0	0			
MIEL2_p	0	0	0	0	0	0	0	0			
SBT52_p2	0	0	0	0	0	0	0	0			
HCD1_p	0	0	0	0	0	0	1	1			
MYB30	0	1	1	1	1	1	1	1			
KCS11_p	0	0	0	0	0	0	0	0			
MYB30	0	0	1	1	1	1	1	1			
sPLA2_a	0	0	0	0	0	0	0	0			
MIEL4_p1	0	0	0	0	0	0	0	0			
CER4_p	0	0	0	0	0	0	1	1			
R_VLCFA	0	0	0	0	0	0	0	1			
MIEL3_p	0	0	0	0	0	0	0	0			
KCS10_p	0	0	0	0	0	0	1	1			
R_VLCFA	0	0	0	0	0	0	0	0	•		

















# Conclusion

- Metabolic network analyses are very insightful to understand cell phenotype.
- Metabolism regulation must be included to get a full picture of cell behaviour.
- These two types of networks can be very large and are of a different nature.
- ► FBA and logical regulatory network modelling provide interesting answers.

